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Not Disclosed by applicants + P212121 ☑ 1RQI ✓ 1R0E Compound Authors Classification Characteristics Authors Classification Characteristics Active Conformation of <u>Farnesyl Pyrophosphate Synthase</u> Bound to Isopentyl Pyrophosphate and <u>Dimethylallyl S-Thiolodiphosphate</u> Glycogen synthase kinase-3 beta in complex with 3-indolyl-4-arylmaleimide inhibitor Release Date: 12-Oct-2004 Exp. Method: X Ray Diffraction Hosfield, D.J., Zhang, Y., Dougan, D.R., Brooun, A., Tari, L.W., Swanson, R.V., Finn, J. Release Date: 02-Mar-2004 Exp. Method: X Ray Diffraction Transferase Resolution: 2.42 Å Allard, J., Nikolchéva, T., Gong, L., Wang, J., Dunten, P., Avnur, Z., Waters, R., Sun, Q., Skinner, B. Mol. Id: 1 Molecule: Glycogen Synthase Kinase 3 Beta Transferase Resolution: 2.25 A Mol. Id: 1 Molecule: Geranyltranstransferasi

The instant
opplication => ✓ 1RQJ ✓ 1RYX **₹** Compound **Authors** Classification Characteristics Crystal structure of hen serum transferrin in apo- form Active Conformation of Farnesyl Pyrophosphate Synthase Bound to Isopentyl Pyrophosphate and Risedronate **Transferase** Resolution: 1.95 Å Release Date: 02-Mar-2004 Exp. Method: X Ray Diffraction Mol. Id: 1 Molecule: Geranyltranstransferase Hosfield, D.J., Zhang, Y., Dougan, D.R., Brooun, A., Tari, L.W., Swanson, R.V., Finn, J.

✓ 1VDC **₹** Classification Characteristics Compound Characteristics Classification Release Date: 12-Mar-1997 Exp. Method: X Ray Diffraction STRUCTURE OF NADPH DEPENDENT THIOREDOXIN REDUCTASE Release Date: 13-Jul-2004 Exp. Method: X Ray Diffraction Mol. Id: 1 Molecule: Nadph Dependent Thioredoxin Reductase Oxidoreductase Dai, S., Saarinen, M., Ramaswamy, S., Meyer, Y., Jacquot, J.P., Eklund, H. Resolution: 2.50 Å Mol. Id: 1 Molecule: Ovotransfefrin **Metal Transport** Resolution: 3.50 Å Thakurta, P.G., Choudhury, D., Dasgupta, R., Dattagupta, J.K.

P4, 2,2

P43212

P4322













Characteristics

Structure of the mediator MED7/MED21 (Med7/Srb7) subcomplex

Release Date: 22-Feb-2005 Exp. Method: X Ray Diffraction Resolution: 3.00 Å

Classification

Fragment: Residues 102 205 Mol. Id: 2 Molecule: RNA Polymerase Ii Holoenzyme Component Srb7 Wutation: L5M, L119M, L125M **Gene Regulation** Mol. Id: 1 Molecule: RNA Polymerase Ii Mediator Complex Protein Med7

Baumli, S., Hoeppner, S., Cramer, P.

Authors

Compound

1 2 **(**

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PDB ID or keyword () Author SEARCH (Advanced Search

Home Search Results Queries ■ Show Query Details ■ Results (11-13 of 13) ■ Refine this Search Results per Page ■ Download Selected P43 212 ✓ 2B4S ✓ 2AYU ✓ 1ZP6 Classification Characteristics Compound **Authors** Compound **Authors** Classification Characteristics Crystal structure of a complex between PTP1B and the insulin receptor tyrosine kinase The structure of nucleosome assembly protein suggests mechanism for histone binding and shuttling Crystal Structure of Atu3015, a Putative Cytidylate Kinase from Agrobacterium tumefaciens, Northeast Structural Genomics Target Structural Genomics Unknown Function Park, Y.J., Luger, K. Chaperone Resolution: 3.00 Å Release Date: 07-Feb-2006_Exp. Method: X Ray Diffraction Mol. Id: 1 Molecule: NucleoSome Assembly Protein Forouhar, F., Abashídze, M., Vorobiev, S.M., Kuzin, A., Conover, K., Acton, T.B., Montelione, G.T., Hunt, J.F., Tong, L. Mol. Id: 1 Molecule: Hypothetical Protein Atu3015 Resolution: 3.20 Å Release Date: 24-May-2005 Exp. Method: X Ray Diffraction

■ Deselect All

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Compound Characteristics **Authors** Classification Release Date: 15-Nov-2005-Exp. Method: X Ray Diffraction Li, S., Depetris, R.S., Barford, D., Chernoff, J., Hubbard, S.R. Molecule: Insulin Receptor Fragment: Protein Kinase Mol. Id: 1 Molecule: Tyrosine Protein Phosphatase Non Receptor Type 1 Mol. Id: 2 Hydrolase/transferase Resolution: 2.30 Å

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STN search
10/651668
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(unit cell dimensions)

=> d his

L1

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(FILE 'HOME' ENTERED AT 16:25:07 ON 01 MAR 2006)
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FILE 'CAPLUS' ENTERED AT 16:25:39 ON 01 MAR 2006

ALL CITATIONS AVAILABLE IN THE RE FORMAT

1840516 S PROTEIN

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L2
      1805951 S ?CRYSTAL?
       52256 S L1 AND L2
L3
L4
        649 S L3 AND (UNIT CELL DIMENSION#)
L5
        25 S L4 AND 88
         2 S L5 AND 174
L6
        -26-S-L3-AND 14122
L7
L8
        665-S-L3-AND-TETRAGONAL
L9
        26-S-L7-AND-L3-
          0.S.L6-ANDL8
L10
=> d L6 1-2
L6 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2006 ACS on STN
AN 1998:463974 CAPLUS
                                                                                Unit cell dimension of a = b = 174 A
c = 88 A
d = \beta = 90, \quad \gamma = 120c
DN 129:199494
TI An internal affinity-tag for purification and crystallization of
   the siderophore receptor FhuA, integral outer membrane protein
   from Escherichia coli K-12
AU Ferguson, Andrew D.; Breed, Jason; Diederichs, Kay; Welte, Wolfram;
   Coulton, James W.
CS Department of Microbiology and Immunology, McGill University, Montreal, —)
   QC, H3A 2B4, Can-
SO Protein Science (1998), 7(7), 1636-1638
   CODEN: PRCIEI; ISSN: 0961-8368
PB Cambridge University Press
DT Journal
LA English
RE.CNT 32 THERE ARE 32 CITED REFERENCES AVAILABLE FOR THIS RECORD
        ALL CITATIONS AVAILABLE IN THE RE FORMAT
L6 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2006 ACS on STN
AN 1998:224997 CAPLUS
                                                                                 a=b=111.0 \text{ Å}
c=174.5 \text{ Å}
x=\beta=90
y=120
DN 128:318765
TI Crystallization of NAD+-dependent phenylalanine dehydrogenase
   from Nocardia sp239
AU Pasquo, A.; Britton, K. L.; Baker, P. J.; Brearley, G.; Hinton, R. J.;
   Moir, A. J. G.; Stillman, T. J.; Rice, D. W.
CS Krebs Institute for Biomolecular Research, Department of Molecular Biology
   and Biotechnology, University of Sheffield, Sheffield, S10 2TN, UK-
SO Acta Crystallographica, Section D. Biological Crystallography (1998),
   D54(2), 269-272
                                                              guar 1
   CODEN: ABCRE6: ISSN: 0907-4449
PB Munksgaard International Publishers Ltd.
DT Journal
LA English
RE.CNT 20
             THERE ARE 20 CITED REFERENCES AVAILABLE FOR THIS RECORD
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